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NiceZyme View of ENZYME: EC 6.3.1.2

Official Name

Glutamate--ammonia ligase.

Alternative Name(s)

Glutamine synthetase.

Reaction catalysed

ATP + L-glutamate + NH(3) <=> ADP + phosphate + L-glutamine

Comment(s)

Also acts, more slowly, on 4-methylene-L-glutamate (cf. EC 6.3.1.7).

Cross-references

Biochemical Pathways; map number(s)

G7

PROSITE

PDOC00162

BRENDA

6.3.1.2

PUMA2

6.3.1.2

PRIAM enzyme-specific profiles

6.3.1.2

Kyoto University LIGAND chemical database

6.3.1.2

IUBMB Enzyme Nomenclature

6.3.1.2

IntEnz

6.3.1.2

MEDLINE

Find literature relating to 6.3.1.2

MetaCyc

6.3.1.2

Q56WN1, GLN11_ARATH;	P14656, GLN11_ORYSA;	Q8LCE1, GLN12_ARATH;
P14654, GLN12_ORYSA;	Q9LVI8, GLN13_ARATH;	Q4W8D0, GLN13_ORYSA;
Q9FMD9, GLN14_ARATH;	Q8GXW5, GLN15_ARATH;	O04867, GLNA1_ALNGL;
P05457, GLNA1_BRAJA;	Q42688, GLNA1_CHLRE;	O22504, GLNA1_DAUCA;
P20477, GLNA1_DROME;	P46033, GLNA1_FRAAL;	Q42899, GLNA1_LOTJA;
P38559, GLNA1_MAIZE;	P04078, GLNA1_MEDSA;	P0A591, GLNA1_MYCBO;
P0A590, GLNA1_MYCTU;	P08282, GLNA1_PEA;	P04770, GLNA1_PHAVU;
P09826, GLNA1_RHILV;	Q59747, GLNA1_RHIME;	P24099, GLNA1_SOYBN;
P77958, GLNA1_STRFL;	Q05542, GLNA1_STRVR;	P51118, GLNA1_VITVI;
Q43127, GLNA2_ARATH;	P04772, GLNA2_BRAJA;	Q42689, GLNA2_CHLRE;
O22506, GLNA2_DAUCA;	P20478, GLNA2_DROME;	P81643, GLNA2_EMIHU;
P20805, GLNA2_FRAAL;	P13564, GLNA2_HORVU;	P38560, GLNA2_MAIZE;
Q9XQ94, GLNA2_MEDSA;	P64246, GLNA2_MYCBO;	P64245, GLNA2_MYCTU;
P14655, GLNA2_ORYSA;	P08281, GLNA2_PEA;	P04771, GLNA2_PHAVU;
P81107, GLNA2_PINPS;	Q02154, GLNA2_RHILP;	P45626, GLNA2_RHIME;
O82560, GLNA2_SOYBN;	P22878, GLNA2_STRHY;	P19432, GLNA2_STRVR;
P51119, GLNA2_VITVI;	Q06378, GLNA3_HORVU;	P14636, GLNA3_LUPAN;
P38561, GLNA3_MAIZE;	Q43785, GLNA3_MEDSA;	P07694, GLNA3_PEA;
P00965, GLNA3_PHAVU;	P31592, GLNA3_RHILP;	O87393, GLNA3_RHIME;

UniProtKB/Swiss-Prot

P38562, GLNA4_MAIZE;	Q43066, GLNA4_PEA;	P15102, GLNA4_PHAVU;
P38563, GLNA5_MAIZE;	Q42624, GLNAC_BRANA;	P25462, GLNAC_MAIZE;
Q9QY94, GLNA_ACOCA;	O00088, GLNA_AGABI;	Q8X169, GLNA_AMAMU;
P00964, GLNA_ANASP;	O66514, GLNA_AQUAE;	O29313, GLNA_ARCFU;
Q75BT9, GLNA_ASHGO;	P10583, GLNA_AZOBR;	P94126, GLNA_AZOCA;
P22248, GLNA_AZOVI;	P19064, GLNA_BACCE;	P15623, GLNA_BACFR;
P12425, GLNA_BACSU;	P15103, GLNA_BOVIN;	Q05650, GLNA_BUTFI;
P34497, GLNA_CAEL;	Q8H2M5, GLNA_CANFA;	Q6FMT6, GLNA_CANGA;
P16580, GLNA_CHICK;	P10656, GLNA_CLOSA;	Q12613, GLNA_COLGL;
P04773, GLNA_CRIGR;	Q96UG9, GLNA_CRYNE;	Q6B4U7, GLNA_DEBHA;
P11600, GLNA_DUNSA;	P0A9C7, GLNA_ECO57;	P0A9C6, GLNA_ECOL6;
P0A9C5, GLNA_ECOLI;	Q96V52, GLNA_EMENI;	P33035, GLNA_FREDI;
Q9UUN6, GLNA_FUSSH;	Q9C2U9, GLNA_GIBFU;	P43794, GLNA_HAEIN;
Q9HNI2, GLNA_HALSA;	P43386, GLNA_HALVO;	Q96UV5, GLNA_HEBCY;
Q9ZLW5, GLNA_HELPJ;	P94845, GLNA_HELPY;	P15104, GLNA_HUMAN;
Q874T6, GLNA_KLULA;	P45627, GLNA_LACDE;	Q9CDL9, GLNA_LACLA;
P23712, GLNA_LACSA;	P52782, GLNA_LUPLU;	Q4R7U3, GLNA_MACFA;
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O27612, GLNA_METTH;	P21154, GLNA_METVO;	P15105, GLNA_MOUSE;
P25821, GLNA_NEIGO;	Q86ZF9, GLNA_NEUCR;	P12424, GLNA_NICPL;
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P09606, GLNA_RAT;	P13499, GLNA_RHOCA;	P43518, GLNA_RHOSH;
P0A1P7, GLNA_SALTI;	P0A1P6, GLNA_SALTY;	Q09179, GLNA_SCHPO;
P0A9C8, GLNA_SHIFL;	P41320, GLNA_SQUAC;	Q5HGC3, GLNA_STAAC;
P60890, GLNA_STAAM;	P99095, GLNA_STAN;	Q6GHC6, GLNA_STAAR;
Q6G9Q4, GLNA_STAAS;	P0A040, GLNA_STAAU;	P0A039, GLNA_STAAW;
Q5HPN2, GLNA_STAEQ;	Q8CSR8, GLNA_STAES;	P15106, GLNA_STRCO;
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P28605, GLNA_SYNP2;	P77961, GLNA_SYNY3;	P36205, GLNA_THEMA;
P07804, GLNA_THIFE;	P51120, GLNA_TRITH;	Q86ZU6, GLNA_TUBBO;
P19904, GLNA_VIBAL;	Q9KNJ2, GLNA_VIBCH;	P32289, GLNA_VIGAC;
P51121, GLNA_XENLA;	Q6C3E0, GLNA_YARLI;	P32288, GLNA_YEAST;

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NiceZyme View of ENZYME: EC 3.5.1.2

Official Name

Glutaminase.

Alternative Name(s)

L-glutamine amidohydrolase.

Reaction catalysed

L-glutamine + H(2)O <=> L-glutamate + NH(3)

Cross-references

Biochemical Pathways; map number(s)

G7

PROSITE

PDOC00132

BRENDA

3.5.1.2

PUMA2

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MEDLINE

Find literature relating to 3.5.1.2

MetaCyc

3.5.1.2

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Q19013, GLS1_CAEEL; Q93650, GLS2_CAEEL; Q81YY0, GLSA1_BACAN;
Q81IB3, GLSA1_BACCR; Q9K9L8, GLSA1_BACHD; O31465, GLSA1_BACSU;
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Q8FK76, GLSA1_ECOL6; P77454, GLSA1_ECOLI; Q83SE1, GLSA1_SHIFL;
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<input type="checkbox"/>	L4	l1 and (method\$ or synthes\$4 or product\$4)	102
<input type="checkbox"/>	L3	l1 and (nakamura or akiyama).in.	3
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<input type="checkbox"/>	L1	glutaminas\$4 same glutamine\$4 same (synthetas\$4 or synthas\$4)	102

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L1 QUE GLUTAMINAS? AND GLUTAMINE?

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FILE 'CAPLUS, BIOSIS, MEDLINE, EMBASE, SCISEARCH, TOXCENTER, PASCAL, USPATFULL, ESBIODBASE, BIOTECHNO, LIFESCI' ENTERED AT 16:33:51 ON 28 AUG

2006

- L2 1535 SEA GLUTAMINAS? AND GLUTAMINE? AND (CORYNEFOR? OR GLUTAMICUM?
OR COLI? OR BACTER? OR BREVIBACT?)
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- L4 342 SEA GLUTAMINAS?(S)(GLUTAMINE?)(S)(CORYNEFOR? OR GLUTAMICUM? OR
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- L5 211 SEA L4(S)(METHOD? OR SYNTH? OR PRODUCT?)
- L6 122 DUP REM L5 (89 DUPLICATES REMOVED)
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L1 QUE GLUTAMINAS? AND GLUTAMINE?

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=> s glutaminas? and glutamine? and (corynefor? or glutamicum? or coli? or bacter?
or brevibact?)

9 FILES SEARCHED...

L2 1535 GLUTAMINAS? AND GLUTAMINE? AND (CORYNEFOR? OR GLUTAMICUM? OR
COLI? OR BACTER? OR BREVIBACT?)

=> s l2 and (method? or product? or synthe?)

7 FILES SEARCHED...

L3 1174 L2 AND (METHOD? OR PRODUCT? OR SYNTHE?)

=> s glutaminas?(s) (glutamine?) (s) (corynefor? or glutamicum? or coli? or bacter? or
brevibact?)

9 FILES SEARCHED...

L4 342 GLUTAMINAS? (S) (GLUTAMINE?) (S) (CORYNEFOR? OR GLUTAMICUM? OR COLI?
OR BACTER? OR BREVIBACT?)

=> s l4(s) (method? or synthe? or product?)

7 FILES SEARCHED...

L5 211 L4(S) (METHOD? OR SYNTHE? OR PRODUCT?)

=> dup rem l5

PROCESSING COMPLETED FOR L5

L6 122 DUP REM L5 (89 DUPLICATES REMOVED)

=> d ti l6 1-122

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TI Active carbohydrate containing protecting reagents for chemical

modifications, their production and use

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TI Use of phosphoketolase for producing useful metabolites
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TI Corynebacterium glutamicum genes encoding metabolic pathway proteins
- L6 ANSWER 9 OF 122 USPATFULL on STN
TI Flea head, nerve cord, hindgut and malpighian tubule nucleic acid molecules, proteins and uses thereof
- L6 ANSWER 10 OF 122 USPATFULL on STN
TI Nucleic acid and amino acid sequences relating to streptococcus pneumoniae for diagnostics and therapeutics
- L6 ANSWER 11 OF 122 USPATFULL on STN
TI Molecular control of transgene segregation and its escape by a recoverable block of function (RBF) system
- L6 ANSWER 12 OF 122 Elsevier BIOBASE COPYRIGHT 2006 Elsevier Science B.V.
on STN DUPLICATE
TI On the two components of pyridoxal 5'-phosphate synthase from Bacillus subtilis
- L6 ANSWER 13 OF 122 Elsevier BIOBASE COPYRIGHT 2006 Elsevier Science B.V.
on STN DUPLICATE
TI Gln-tRNA^{sup.G.sup.l.sup.n} formation from Glu-tRNA^{sup.G.sup.l.sup.n} requires cooperation of an asparaginase and a Glu-tRNA^{sup.G.sup.l.sup.n} kinase
- L6 ANSWER 14 OF 122 Elsevier BIOBASE COPYRIGHT 2006 Elsevier Science B.V.
on STN DUPLICATE
TI Analysis of the vitamin B6 biosynthesis pathway in the human malaria parasite Plasmodium falciparum
- L6 ANSWER 15 OF 122 CAPLUS COPYRIGHT 2006 ACS on STN DUPLICATE 6
TI Characterization of LtsA from Rhodococcus erythropolis, an enzyme with glutamine amidotransferase activity
- L6 ANSWER 16 OF 122 Elsevier BIOBASE COPYRIGHT 2006 Elsevier Science B.V.
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TI Lid L11 of the glutamine amidotransferase domain of CTP synthase mediates allosteric GTP activation of glutaminase activity

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on STN DUPLICATE 8
TIEN Characterization of salt-tolerant glutaminase from *Stenotrophomonas maltophilia* NYW-81 and its application in Japanese soy sauce fermentation

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on STN
TI Structure-function studies on the iron-sulfur flavoenzyme glutamate synthase: An unexpectedly complex self-regulated enzyme

L6 ANSWER 19 OF 122 CAPLUS COPYRIGHT 2006 ACS on STN
TI Fermentative production of L-glutamine by genetically modified *Corynebacterium glutamicum*

L6 ANSWER 20 OF 122 USPATFULL on STN
TI Method for producing L-glutamine and L-glutamine producing bacterium

L6 ANSWER 21 OF 122 USPATFULL on STN
TI Compounds for the modulation of the glycolysis enzyme and/or transaminase complex

L6 ANSWER 22 OF 122 USPATFULL on STN
TI *Staphylococcus aureus* polynucleotides and sequences

L6 ANSWER 23 OF 122 USPATFULL on STN
TI Molecular control of transgene segregation and its escape by a recoverable block of function (rbf) system

L6 ANSWER 24 OF 122 USPATFULL on STN
TI Nucleic acid and amino acid sequences relating to *Streptococcus pneumoniae* for diagnostics and therapeutics

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on STN DUPLICATE
TI Domain organization of *Salmonella typhimurium* formylglycinamide ribonucleotide amidotransferase revealed by X-ray crystallography

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TI Long-range allosteric transitions in carbamoyl phosphate synthetase

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TI Characterization of the products of the genes SNO1 and SNZ1 involved in pyridoxine synthesis in *Saccharomyces cerevisiae*

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on STN
TI Molecular cloning, overexpression, and purification of *Micrococcus luteus* K-3-type glutaminase from *Aspergillus oryzae* RIB40

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on STN DUPLICATE
TI Inhibition of *E. coli* CTP synthase by the "positive" allosteric effector GTP

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on STN
TI Regulation of transcription and activity of *Rhizobium etli* glutaminase a

L6 ANSWER 31 OF 122 USPATFULL on STN
TI Microbial culture with enhanced glutaminase activity and utilization thereof

L6 ANSWER 32 OF 122 USPATFULL on STN
 TI Identification of modulatory molecules using inducible promoters

L6 ANSWER 33 OF 122 USPATFULL on STN
 TI Flea head, nerve cord, hindgut and malpighian tubule nucleic acid molecules, proteins and uses thereof

L6 ANSWER 34 OF 122 USPATFULL on STN
 TI STAPHYLOCOCCUS AUREUS POLYNUCLEOTIDES AND SEQUENCES

L6 ANSWER 35 OF 122 USPATFULL on STN
 TI Nucleic acid and amino acid sequences relating to Enterococcus faecalis for diagnostics and therapeutics

L6 ANSWER 36 OF 122 USPATFULL on STN
 TI Staphylococcus aureus polynucleotides and sequences

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 TI Gain of glutaminase function in mutants of the ammonia-specific frog carbamoyl phosphate synthetase

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 TIEN Functional characterization of a salt- and thermotolerant glutaminase from Lactobacillus rhamnosus

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 TIEN Microbial glutaminase: biochemistry, molecular approaches and applications in the food industry
 Enzyme biochemistry and biotechnology. A collection of papers dedicated to Professor Dr. Kenji Soda in honor of his 70th birthday

L6 ANSWER 40 OF 122 Elsevier BIOBASE COPYRIGHT 2006 Elsevier Science B.V. on STN
 TI Revisiting the steady state kinetic mechanism of glutamine-dependent asparagine synthetase from Escherichia coli

L6 ANSWER 41 OF 122 USPATFULL on STN
 TI Utilization of Wolinella succinogenes asparaginase to treat diseases associated with asparagine dependence

L6 ANSWER 42 OF 122 USPATFULL on STN
 TI STREPTOCOCCUS PNEUMONIAE POLYNUCLEOTIDES AND SEQUENCES

L6 ANSWER 43 OF 122 CAPLUS COPYRIGHT 2006 ACS on STN DUPLICATE 16
 TI Steady-state kinetics of the glutaminase reaction of CTP synthase from Lactococcus lactis. The role of the allosteric activator GTP in coupling between glutamine hydrolysis and CTP synthesis

L6 ANSWER 44 OF 122 USPATFULL on STN
 TI Utilization of Wolinella succinogenes asparaginase to treat diseases associated with asparagine dependence

L6 ANSWER 45 OF 122 BIOTECHNO COPYRIGHT 2006 Elsevier Science B.V. on STN DUPLICATE
 TI A Novel Carbamoyl-Phosphate Synthetase from Aquifex aeolicus

L6 ANSWER 46 OF 122 Elsevier BIOBASE COPYRIGHT 2006 Elsevier Science B.V. on STN DUPLICATE
 TI Imidazole glycerol phosphate synthase from Thermotoga maritima. Quaternary structure, steady-state kinetics, and reaction mechanism of the bienzyme complex

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on STN DUPLICATE
TI Structural basis for the activity and substrate specificity of *Erwinia chrysanthemi* L-asparaginase

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TI Protein-glutaminase from *Chryseobacterium proteolyticum*, an enzyme that deamidates glutamyl residues in proteins: Purification, characterization and gene cloning

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TI Mechanism for acivicin inactivation of triad glutamine amidotransferases

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TI Characterization of Glutaminase from Triticale

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on STN
TIEN Carbamoyl-phosphate synthetases (CPS) in lactic acid bacteria and other Gram-positive bacteria
TIFR Les bacteries lactiques : du cognitif a l'application

L6 ANSWER 52 OF 122 USPATFULL on STN
TI Methods for therapy of cancer

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TI Temperature-dependent function of the glutamine phosphoribosylpyrophosphate amidotransferase ammonia channel and coupling with glycylamide ribonucleotide synthetase in a hyperthermophile

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on STN DUPLICATE
TI An engineered blockage within the ammonia tunnel of carbamoyl phosphate synthetase prevents the use of glutamine as a substrate but not ammonia

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on STN DUPLICATE
TI Expression and purification of imidazole glycerol phosphate synthase from *Saccharomyces cerevisiae*

L6 ANSWER 56 OF 122 USPATFULL on STN
TI Pharmaceutical and diet formulations for the prophylaxis and treatment of gastrointestinal disorders

L6 ANSWER 57 OF 122 USPATFULL on STN
TI Compositions and methods for treating and preventing pathologies including cancer

L6 ANSWER 58 OF 122 USPATFULL on STN
TI Compositions and methods for therapy and prevention of cancer, AIDS, and anemia

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TI Functional linkage between the glutaminase and synthetase domains of carbamoyl-phosphate synthetase - Role of serine 44 in carbamoyl-phosphate synthetase-aspartate carbamoyltransferase-dihydroorotase (CAD)

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on STN DUPLICATE
TI Deconstruction of the catalytic array within the amidotransferase subunit of carbamoyl phosphate synthetase

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 TI Poly(ethylene glycol)-bovine serum albumin hydrogel as a matrix for
 enzyme immobilization. In vitro biochemical characterization

L6 ANSWER 62 OF 122 USPATFULL on STN
 TI Compositions and methods for treating and preventing pathologies
 including cancer

L6 ANSWER 63 OF 122 USPATFULL on STN
 TI Aptamers specific for biomolecules and methods of making

L6 ANSWER 64 OF 122 USPATFULL on STN
 TI Methods of inducing the production of hemoglobin and treating
 pathologies associated with abnormal hemoglobin activity using
 phenylacetic acids and derivatives thereof

L6 ANSWER 65 OF 122 USPATFULL on STN
 TI Compositions and methods for therapy and prevention of pathologies
 including cancer, AIDS, and anemia

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 TI Methods for promoting wound healing

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 on STN DUPLICATE
 TI The recombinant α subunit of glutamate synthase: Spectroscopic and
 catalytic properties

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 TI Carbamoyl-phosphate synthetase II in kinetoplastids

L6 ANSWER 69 OF 122 USPATFULL on STN
 TI Method of stabilizing enzyme conjugates

L6 ANSWER 70 OF 122 USPATFULL on STN
 TI Methods for treating neoplastic conditions using phenylacetic acid and
 derivatives thereof

L6 ANSWER 71 OF 122 USPATFULL on STN
 TI Methods for prevention of cancer using phenylacetic acids and
 derivatives thereof

L6 ANSWER 72 OF 122 USPATFULL on STN
 TI Methods for inducing differentiation of a cell using phenylacetic acid
 and derivatives

L6 ANSWER 73 OF 122 USPATFULL on STN
 TI Compositions and methods for therapy and prevention of pathologies
 including cancer, AIDS and anemia

L6 ANSWER 74 OF 122 USPATFULL on STN
 TI Compositions and methods for treating and preventing pathologies
 including cancer

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 DUPLICATE
 TI The smallest carbamoyl-phosphate synthetase. A single catalytic subdomain
 catalyzes all three partial reactions

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 TI Activation by fusion of the glutaminase and synthetase subunits of
 Escherichia coli carbamoyl-phosphate synthetase

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TI Trapping an activated conformation of mammalian carbamyl-phosphate
synthetase

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TI Catalytic activity of the N-terminal domain of Escherichia coli
asparagine synthetase B can be reengineered by single-point mutation

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TI Low glutamine concentrations induce phenotypical and functional
differentiation of U937 myelomonocytic cells

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TI In vivo mutational analysis of highly conserved amino acid residues of
the small subunit Cpalp of the Carbamylphosphate synthetase of
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TIEN The use of glutamine in the treatment of gastrointestinal disorders in
man

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TI Structure and function of the glutamine phosphoribosylpyrophosphate
amidotransferase glutamine site and communication with the
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TI Probing the mechanism of nitrogen transfer in Escherichia coli asparagine
synthetase by using heavy atom isotope effects

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TIEN The role of glutaminase in Rhizobium etli : studies with a new mutant

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on STN DUPLICATE 35
TI SUBSTRUCTURE OF THE AMIDOTRANSFERASE DOMAIN OF MAMMALIAN
CARBAMYL-PHOSPHATE SYNTHETASE

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TIEN Effect of germfree state on the capacities of isolated rat colonocytes to
metabolize n-butyrate, glucose, and glutamine

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TIEN Role of the glutamine transaminase- ω -amidase pathway and
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TI Mapping the structural domains of E. coli carbamoyl phosphate synthetase
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TI Structural characterization of Pseudomonas 7A glutaminase-asparaginase

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TIEN Glutamine-dependent nitrogen transfer in Escherichia coli asparagine synthetase B : searching for the catalytic triad

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TI A molecular wedge for triggering the amidotransferase activity of carbamoyl phosphate synthetase

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TI Conjugates of monophenyl thyroid analogs useful in assays

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TI Imidazole glycerol phosphate synthase: The glutamine amidotransferase in histidine biosynthesis.

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TI Substitution of Glu841 by lysine in the carbamate domain of carbamyl phosphate synthetase alters the catalytic properties of the glutaminase subunit

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TIEN Endotoxin stimulates lymphocytes glutaminase expression

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TI A novel method of production of theanine by immobilized Pseudomonas nitroreducens cells

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on STN DUPLICATE 43

TIEN A continuous production method for theanine by immobilized Pseudomonas nitroreducens cells

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TIEN Dexamethasone stimulation of glutaminase expression in mesenteric lymph nodes. Discussion

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TIEN Extracellular L-glutaminase production by marine bacteria

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TIEN Detection of an enzyme bound γ -glutamyl acyl ester of carbamyl phosphate synthetase of Escherichia coli

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TI Mechanistic studies of glutaminase activity of a glutamine amidotransferase, carbamoyl phosphate synthetase from Escherichia coli

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TI Role of the four conserved histidine residues in the amidotransferase domain of carbamoyl phosphate synthetase

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TI The catalytic mechanism of the amidotransferase domain of the syrian hamster multifunctional protein CAD: Evidence for a CAD-glutamyl covalent intermediate in the formation of carbamyl phosphate

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TI ENDOTOXIN AND RENAL GLUTAMINE METABOLISM.

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TI Mammalian carbamyl phosphate synthetase (CPS). cDNA sequence and evolution of the CPS domain of the Syrian hamster multifunctional protein CAD

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TI Escherichia coli carbamoyl phosphate-synthetase: Domains of glutaminase and synthetase-subunit interaction

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TI Formylglycinamide ribonucleotide synthetase from Escherichia coli: Cloning, sequencing, overproduction, isolation, and characterization

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TI Formation of gamma -glutamyl peptides by glutaminase of Aspergillus oryzae .

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TIEN A mathematical model for the growth of a single cell of E. coli on a glucose/glutamine/ammonium medium

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TIEN L-asparaginase effects on inhibition of protein synthesis and lowering of the glutamine content in cultured rat hepatocytes

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TI The gene coding for carbamoyl-phosphate synthetase I was formed by fusion of an ancestral glutaminase gene and a synthetase gene

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TI L-Glutamine and L-glutamate. UV-method with glutaminase and glutamate dehydrogenase

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TI Inhibition of glucosamine-6-phosphate synthetase from bacteria by anticapsin

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TI Characterization of the effects of asparaginase from Escherichia coli and a glutaminase-free asparaginase from vibrio Succinogenes on specific cell-mediated cytotoxicity

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TI MECHANISM OF SENSITIVITY OF CULTURED PANCREATIC CARCINOMA TO ASPARAGINASE.

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TI 'Regulation of glutaminase B in Escherichia coli. III. Control by nucleotides and divalent cations.

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TI PROPERTIES OF ANTHRANILATE SYNTHETASE COMPONENT II FROM
PSEUDOMONAS-PUTIDA.

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TI Inhibition by dithiothreitol of the utilization of glutamine by carbamyl
phosphate synthetase. Evidence for formation of hydrogen peroxide

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TI Adenosine 3',5'-cyclic monophosphate control of the enzymes of glutamine
metabolism in Escherichia coli

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TI ANTI NEOPLASTIC ACTIVITY OF CERTAIN BACTERIAL ENZYME PREPARATIONS.

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TI Bacterial production of glutamic acid in stored comminuted beef

=> d ibib abs 16 8 15 19 20 28 31 39 59 67 95 102 112 122

L6 ANSWER 8 OF 122 USPATFULL on STN

ACCESSION NUMBER: 2005:299042 USPATFULL

TITLE: Corynebacterium glutamicum genes encoding metabolic
pathway proteins

INVENTOR(S): Pompejus, Markus, Freinsheim, GERMANY, FEDERAL REPUBLIC
OF
Kroger, Burkhard, Limburgerhof, GERMANY, FEDERAL
REPUBLIC OF
Schroder, Hartwig, Nussloch, GERMANY, FEDERAL REPUBLIC
OF
Zelder, Oskar, Speyer, GERMANY, FEDERAL REPUBLIC OF
Haberhauer, Gregor, Limburgerhof, GERMANY, FEDERAL
REPUBLIC OF

PATENT ASSIGNEE(S): BASF Aktiengesellschaft, Ludwigshafen, GERMANY, FEDERAL
REPUBLIC OF (non-U.S. corporation)

	NUMBER	KIND	DATE
PATENT INFORMATION:	US 2005260707	A1	20051124
APPLICATION INFO.:	US 2005-55822	A1	20050211 (11)
RELATED APPLN. INFO.:	Continuation of Ser. No. US 2000-606740, filed on 23 Jun 2000, ABANDONED		

	NUMBER	DATE
PRIORITY INFORMATION:	DE 1999-19932125	19990709
	DE 1999-19932227	19990709
	DE 1999-19932228	19990709
	DE 1999-19932230	19990709
	DE 1999-19933005	19990714
	DE 1999-19933006	19990714
	DE 1999-19940764	19990827
	DE 1999-19940766	19990827
	DE 1999-19940832	19990827
	DE 1999-19941378	19990831
	DE 1999-19941379	19990831
	DE 1999-19942077	19990903
	DE 1999-19942079	19990903
	DE 1999-19931418	19990708
	DE 1999-19932126	19990709
	DE 1999-19932229	19990709
	DE 1999-19941396	19990831
	DE 1999-19942087	19990903

DE 1999-19930476	19990701
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DE 1999-19942088	19990903
DE 1999-19942124	19990903
DE 1999-19932928	19990714
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DE 1999-19942086	19990903
DE 1999-19942095	19990903
DE 1999-19942129	19990903
US 1999-141031P	19990625 (60)
US 1999-142101P	19990702 (60)
US 1999-148613P	19990812 (60)
US 2000-187970P	20000309 (60)

DOCUMENT TYPE: Utility
FILE SEGMENT: APPLICATION
LEGAL REPRESENTATIVE: LAHIVE & COCKFIELD, LLP., 28 STATE STREET, BOSTON, MA, 02109, US
NUMBER OF CLAIMS: 38
EXEMPLARY CLAIM: 1
LINE COUNT: 8777

CAS INDEXING IS AVAILABLE FOR THIS PATENT.

AB Isolated nucleic acid molecules, designated MP nucleic acid molecules, which encode novel MP proteins from *Corynebacterium glutamicum* are described. The invention also provides antisense nucleic acid molecules, recombinant expression vectors containing MP nucleic acid molecules, and host cells into which the expression vectors have been introduced. The invention still further provides isolated MP proteins, mutated MP proteins, fusion proteins, antigenic peptides and methods for the improvement of production of a desired compound from *C. glutamicum* based on genetic engineering of MP genes in this organism.

CAS INDEXING IS AVAILABLE FOR THIS PATENT.

L6 ANSWER 15 OF 122 CAPLUS COPYRIGHT 2006 ACS on STN DUPLICATE 6
ACCESSION NUMBER: 2005:330029 CAPLUS
DOCUMENT NUMBER: 143:22102
TITLE: Characterization of LtsA from *Rhodococcus erythropolis*, an enzyme with glutamine

AUTHOR(S): amidotransferase activity
 Mitani, Yasuo; Meng, Xian Ying; Kamagata, Yoichi;
 Tamura, Tomohiro
 CORPORATE SOURCE: Proteolysis and Protein Turnover Research Group,
 Research Institute of Genome-Based Biofactory,
 National Institute of Advanced Industrial Science and
 Technology (AIST), Toyohira-ku, Japan
 SOURCE: Journal of Bacteriology (2005), 187(8), 2582-2591
 CODEN: JOBAAY; ISSN: 0021-9193
 PUBLISHER: American Society for Microbiology
 DOCUMENT TYPE: Journal
 LANGUAGE: English

AB The nocardioform actinomycete *Rhodococcus erythropolis* has a characteristic cell wall structure. The cell wall is composed of arabinogalactan and mycolic acid and is highly resistant to the cell wall-lytic activity of lysozyme (muramidase). In order to improve the isolation of recombinant proteins from *R. erythropolis* host cells (N. Nakashima and T. Tamura, *Biotechnol. Bioeng.* 86:136-148, 2004), we isolated two mutants, L-65 and L-88, which are susceptible to lysozyme treatment. The lysozyme sensitivity of the mutants was complemented by expression of *Corynebacterium glutamicum* *ltsA*, which codes for an enzyme with glutamine amidotransferase activity that results from coupling of two reactions (a glutaminase activity and a synthetase activity). The lysozyme sensitivity of the mutants was also complemented by *ltsA* homologs from *Bacillus subtilis* and *Mycobacterium tuberculosis*, but the homologs from *Streptomyces coelicolor* and *Escherichia coli* did not complement the sensitivity. This result suggests that only certain *LtsA* homologs can confer lysozyme resistance. Wild-type recombinant *LtsA* from *R. erythropolis* showed glutaminase activity, but the *LtsA* enzymes from the L-88 and L-65 mutants displayed drastically reduced activity. Interestingly, an *ltsA* disruptant mutant, which expressed the mutated *LtsA*, changed from lysozyme sensitive to lysozyme resistant when NH_4Cl was added into the culture media. The glutaminase activity of the *LtsA* mutants inactivated by site-directed mutagenesis was also restored by addition of NH_4Cl , indicating that NH_3 can be used as an amide donor mol. Taken together, these results suggest that *LtsA* is critically involved in mediating lysozyme resistance in *R. erythropolis* cells.

REFERENCE COUNT: 38 THERE ARE 38 CITED REFERENCES AVAILABLE FOR THIS RECORD. ALL CITATIONS AVAILABLE IN THE RE FORMAT

L6 ANSWER 19 OF 122 CAPLUS COPYRIGHT 2006 ACS on STN
 ACCESSION NUMBER: 2004:446945 CAPLUS
 DOCUMENT NUMBER: 141:5878
 TITLE: Fermentative production of L-glutamine by genetically modified *Corynebacterium glutamicum*
 INVENTOR(S): Nakamura, Jun; Akiyama, Kayo
 PATENT ASSIGNEE(S): Ajinomoto Co., Inc., Japan
 SOURCE: Eur. Pat. Appl., 37 pp.
 CODEN: EPXXDW
 DOCUMENT TYPE: Patent
 LANGUAGE: English
 FAMILY ACC. NUM. COUNT: 1
 PATENT INFORMATION:

PATENT NO.	KIND	DATE	APPLICATION NO.	DATE
EP 1424397	A1	20040602	EP 2003-26890	20031124
R: AT, BE, CH, DE, DK, ES, FR, GB, GR, IT, LI, LU, NL, SE, MC, PT, IE, SI, LT, LV, FI, RO, MK, CY, AL, TR, BG, CZ, EE, HU, SK				
US 2004152175	A1	20040805	US 2003-720177	20031125
BR 2003005314	A	20040831	BR 2003-5314	20031125
CN 1502689	A	20040609	CN 2003-10124084	20031126
JP 2004187684	A2	20040708	JP 2003-395175	20031126
PRIORITY APPLN. INFO.:			JP 2002-342287	A 20021126

AB L-Glutamine is produced by culturing a coryneform bacterium having L-glutamine- producing ability and modified so that intracellular glutaminase activity is reduced, and preferably also modified so that intracellular glutamine synthetase activity is enhanced. The method of production includes culturing the bacterium in a medium, followed by accumulation of L-glutamine in the medium and collecting the L-glutamine from the medium.

L6 ANSWER 20 OF 122 USPATFULL on STN

ACCESSION NUMBER: 2004:196870 USPATFULL
TITLE: Method for producing L-glutamine and L-glutamine producing bacterium
INVENTOR(S): Nakamura, Jun, Kawasaki-shi, JAPAN
Akiyama, Kayo, Kawasaki-shi, JAPAN

	NUMBER	KIND	DATE
PATENT INFORMATION:	US 2004152175	A1	20040805
APPLICATION INFO.:	US 2003-720177	A1	20031125 (10)

	NUMBER	DATE
PRIORITY INFORMATION:	JP 2002-342287	20021126
DOCUMENT TYPE:	Utility	
FILE SEGMENT:	APPLICATION	
LEGAL REPRESENTATIVE:	AJINOMOTO CORPORATE SERVICES, LLC, INTELLECTUAL PROPERTY DEPARTMENT, 1120 CONNECTICUT AVE., N.W., WASHINGTON, DC, 20036	
NUMBER OF CLAIMS:	11	
EXEMPLARY CLAIM:	1	
NUMBER OF DRAWINGS:	4 Drawing Page(s)	
LINE COUNT:	1523	

CAS INDEXING IS AVAILABLE FOR THIS PATENT.

AB L-glutamine is produced by culturing a coryneform bacterium having L-glutamine-producing ability and modified so that intracellular glutaminase activity is reduced, and preferably also modified so that intracellular glutamine synthetase activity is enhanced. The method of production includes culturing the bacterium in a medium, followed by accumulation of L-glutamine in the medium and collecting the L-glutamine from the medium.

CAS INDEXING IS AVAILABLE FOR THIS PATENT.

L6 ANSWER 28 OF 122 Elsevier BIOBASE COPYRIGHT 2006 Elsevier Science B.V. on STN

ACCESSION NUMBER: 2004296125 ESBIODASE
TITLE: Molecular cloning, overexpression, and purification of Micrococcus luteus K-3-type glutaminase from Aspergillus oryzae RIB40
AUTHOR: Masuo N.; Ito K.; Yoshimune K.; Hoshino M.; Matsushima K.; Koyama Y.; Moriguchi M.
CORPORATE SOURCE: E-mail: mmorigu@cc.oita-u.ac.jp
SOURCE: Protein Expression and Purification, (2004), 38/2 (272-278), 24 reference(s)
CODEN: PEXPEJ ISSN: 1046-5928
PUBLISHER ITEM IDENT.: S1046592804003080
DOCUMENT TYPE: Journal; Article
COUNTRY: United States
LANGUAGE: English
SUMMARY LANGUAGE: English

AB We have for the first time found and cloned the cDNA (AoglsA) of Aspergillus oryzae RIB40, which encodes a 49.9-kDa protein sharing 40% homology with the salt-tolerant glutaminase of Micrococcus

luteus K-3 (Micrococcus glutaminase). AoglsA was subcloned into a series of expression vectors and expressed in Saccharomyces cerevisiae and Escherichia coli. The gene product, which we named AoGls, showed glutaminase activity and was produced in a cell wall fraction of S. cerevisiae and a soluble protein in E. coli. The highest expression level of 186 U/mg was obtained when the AoglsA was inserted into six bases downstream of the Shine-Dalgarno (SD) sequence of pKK223-3 and expressed in E. coli Rosetta (DE3). AoGls was purified by SuperQ-TOYOPEARL, glutamine affinity chromatography, and Butyl-TOYOPEARL. This is the first report on the overexpression and purification of a M. luteus K-3-type glutaminase cloned from an eucaryote. .COPYRG. 2004 Elsevier Inc. All rights reserved.

L6 ANSWER 31 OF 122 USPATFULL on STN
 ACCESSION NUMBER: 2003:180816 USPATFULL
 TITLE: Microbial culture with enhanced glutaminase activity and utilization thereof
 INVENTOR(S): Yuasa, Ari, Kawasaki-shi, JAPAN
 Okamura, Hideki, Kawasaki-shi, JAPAN
 Kataoka, Jiro, Kawasaki-shi, JAPAN
 PATENT ASSIGNEE(S): AJINOMOTO CO. INC., Tokyo, JAPAN (non-U.S. corporation)

	NUMBER	KIND	DATE
PATENT INFORMATION:	US 2003124646	A1	20030703
APPLICATION INFO.:	US 2002-285642	A1	20021101 (10)
RELATED APPLN. INFO.:	Continuation of Ser. No. US 2000-647923, filed on 7 Dec 2000, ABANDONED A 371 of International Ser. No. WO 1999-JP1983, filed on 14 Apr 1999, UNKNOWN		

	NUMBER	DATE
PRIORITY INFORMATION:	JP 1998-121621	19980416
DOCUMENT TYPE:	Utility	
FILE SEGMENT:	APPLICATION	
LEGAL REPRESENTATIVE:	OBLON, SPIVAK, MCCLELLAND, MAIER & NEUSTADT, P.C., 1940 DUKE STREET, ALEXANDRIA, VA, 22314	
NUMBER OF CLAIMS:	8	
EXEMPLARY CLAIM:	1	
LINE COUNT:	720	

CAS INDEXING IS AVAILABLE FOR THIS PATENT.

AB A microbial culture having an increased glutaminase activity is produced by releasing catabolite repression of said glutaminase during incubation of a microorganism capable of producing glutaminase, and feeding a nitrogen source in the intermediate stage of the incubation as required.

Protein is subjected to a reaction with the thus prepared microbial culture in the presence of proteolytic enzymes and either in the absence of sodium chloride or in the presence of sodium chloride at a concentration of 3% (weight/volume) or less, thereby giving hydrolyzed protein which has a potent flavoring effect and is highly useful as a food seasoning.

CAS INDEXING IS AVAILABLE FOR THIS PATENT.

L6 ANSWER 39 OF 122 PASCAL COPYRIGHT 2006 INIST-CNRS. ALL RIGHTS RESERVED.
 on STN DUPLICATE 15
 ACCESSION NUMBER: 2004-0026662 PASCAL
 COPYRIGHT NOTICE: Copyright .COPYRG. 2004 INIST-CNRS. All rights reserved.
 TITLE (IN ENGLISH): Microbial glutaminase: biochemistry, molecular approaches and applications in the food industry
 Enzyme biochemistry and biotechnology. A collection of papers dedicated to Professor Dr. Kenji Soda in honor

of his 70th birthday

AUTHOR: NANDAKUMAR Renu; YOSHIMUNE Kazuaki; WAKAYAMA Mamoru;
MORIGUCHI Mitsuaki
NAKAJIMA Nobuyoshi (ed.)

CORPORATE SOURCE: Department of Chemical and Biochemical Engineering,
University of Maryland Baltimore County, 1000 Hilltop
Circle, Baltimore, MD 21250, United States; Department
of Applied Chemistry, Faculty of Engineering, Oita
University, Dannoharu 700, Oita 870-1192, Japan;
Department of Bioscience and Biotechnology, Faculty of
Science and Engineering, Ritsumeikan University, Noji,
Kusatsu, Shiga 525-8577, Japan
Department of Nutritional Science, Faculty of Health
and Welfare Science, Okayama Prefectural University,
Soja, 719-1197 Okayama, Japan

SOURCE: Journal of molecular catalysis. B, Enzymatic, (2003),
23(2-6), 87-100, 77 refs.
ISSN: 1381-1177

DOCUMENT TYPE: Journal

BIBLIOGRAPHIC LEVEL: Analytic

COUNTRY: Netherlands

LANGUAGE: English

AVAILABILITY: INIST-17107B, 354000112883890030

AN 2004-0026662 PASCAL

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AB Glutaminase is widely distributed in microorganisms including
bacteria, yeast and fungi. The enzyme mainly catalyzes the
hydrolysis of γ -amido bond of L- glutamine. In addition,
some enzymes also catalyze γ -glutamyl transfer reaction. A highly
savory amino acid, L-glutamic acid and a taste-enhancing amino acid of
infused green tea, theanine can be synthesized by employing
hydrolytic or transfer reaction catalyzed by glutaminase.
Therefore, glutaminase is one of the most important
flavor-enhancing enzymes in food industries. In this review, subsequent
to a discussion on the definition of glutaminase, the enzymatic
properties, applications of glutaminase in the food industry,
and occurrence and distribution of the enzyme are described. We then
illustrate the gene cloning, primary structure, and 3D-structure of
glutaminase. Finally, to facilitate the future applications of
glutaminase in food fermentations, the mechanisms of action of
salt-tolerant glutaminase are briefly discussed.

L6 ANSWER 59 OF 122 SCISEARCH COPYRIGHT (c) 2006 The Thomson Corporation
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ACCESSION NUMBER: 1999:759192 SCISEARCH

THE GENUINE ARTICLE: 241ZA

TITLE: Functional linkage between the glutaminase and synthetase
domains of carbamoyl-phosphate synthetase - Role of serine
44 in carbamoyl-phosphate synthetase-aspartate
carbamoyltransferase-dihydroorotase (CAD)

AUTHOR: Hewagama A; Guy H I; Vickrey J F; Evans D R (Reprint)

CORPORATE SOURCE: Wayne State Univ, Sch Med, Dept Biochem & Mol Biol,
Detroit, MI 48201 USA (Reprint)

COUNTRY OF AUTHOR: USA

SOURCE: JOURNAL OF BIOLOGICAL CHEMISTRY, (1 OCT 1999) Vol. 274,
No. 40, pp. 28240-28245.
ISSN: 0021-9258.

PUBLISHER: AMER SOC BIOCHEMISTRY MOLECULAR BIOLOGY INC, 9650
ROCKVILLE PIKE, BETHESDA, MD 20814 USA.

DOCUMENT TYPE: Article; Journal

LANGUAGE: English

REFERENCE COUNT: 50

ENTRY DATE: Entered STN: 1999
Last Updated on STN: 1999

ABSTRACT IS AVAILABLE IN THE ALL AND IALL FORMATS

AB

Mammalian carbamoyl-phosphate synthetase is part of carbamoyl-phosphate synthetase-aspartate carbamoyltransferase-dihydroorotase (CAD), a multifunctional protein that also catalyzes the second and third steps of pyrimidine biosynthesis. Carbamoyl phosphate synthesis requires the concerted action of the glutaminase (GLN) and carbamoyl-phosphate synthetase domains of CAD. There is a functional linkage between these domains such that glutamine hydrolysis on the GLN domain does not occur at a significant rate unless ATP and HCO_3^- , the other substrates needed for carbamoyl phosphate synthesis, bind to the synthetase domain. The GLN domain consists of catalytic and attenuation subdomains. In the separately cloned GLN domain, the catalytic subdomain is down-regulated by interactions with the attenuation domain, a process thought to be part of the functional linkage. Replacement of Ser(44) in the GLN attenuation domain with alanine increases the $k(\text{cat})/K_m$ for glutamine hydrolysis 680-fold. The formation of a functional hybrid between the mammalian Ser(44) GLN domain and the *Escherichia coli* carbamoyl-phosphate synthetase large subunit had little effect on glutamine hydrolysis. In contrast, ATP and HCO_3^- did not stimulate the glutaminase activity, indicating that the interdomain linkage had been disrupted. In accord with this interpretation, the rate of glutamine hydrolysis and carbamoyl phosphate synthesis were no longer coordinated. Approximately 3 times more glutamine was hydrolyzed by the Ser(44) --> Ala mutant than that needed for carbamoyl phosphate synthesis. Ser(44), the only attenuation subdomain residue that extends into the GLN active site, appears to be an integral component of the regulatory circuit that phases glutamine hydrolysis and carbamoyl phosphate synthesis.

L6 ANSWER 67 OF 122 Elsevier BIOBASE COPYRIGHT 2006 Elsevier Science B.V.
on STN DUPLICATE

ACCESSION NUMBER: 1998047215 ESBIOBASE
TITLE: The recombinant α subunit of glutamate synthase: Spectroscopic and catalytic properties
AUTHOR: Vanoni M.A.; Fischer F.; Ravasio S.; Verzotti E.; Edmondson D.E.; Hagen W.R.; Zanetti G.; Curti B.
CORPORATE SOURCE: M.A. Vanoni, Dipto. di Fisiol./Biochim. Generali, Universita degli Studi di Milano, Via Celoria 26, 20133 Milano, Italy.
E-mail: mav@imiucca.csi.unimi.it
SOURCE: Biochemistry, (17 FEB 1998), 37/7 (1828-1838), 28 reference(s)
CODEN: BICHAW ISSN: 0006-2960
DOCUMENT TYPE: Journal; Article
COUNTRY: United States
LANGUAGE: English
SUMMARY LANGUAGE: English

AB As part of our studies of *Azospirillum brasilense* glutamate synthase, a complex iron-sulfur flavoprotein, we have overproduced the two enzyme subunits separately in *Escherichia coli*. The β subunit (53.2 kDa) was demonstrated to contain the site of NADPH oxidation of glutamate synthase and the FAD cofactor, which was identified as Flavin 1 of glutamate synthase, the flavin located at the site of NADPH oxidation. We now report the overproduction of the glutamate synthase α subunit (162 kDa), which is purified to homogeneity in a stable form. This subunit contains FMN as the ravin cofactor which exhibits the properties of Flavin 2 of glutamate synthase: reactivity with sulfite to yield a flavin-N(5)-sulfite addition product ($K(d) = 2.6 \pm 0.22 \text{ mM}$), lack of reactivity with NADPH, reduction by L-glutamate, and reoxidation by 2-oxoglutarate and glutamine. Thus, FMN is the ravin located at the site of reduction of the iminoglutarate formed on the addition of glutamine amide group to the C(2) carbon of 2-oxoglutarate. The glutamate synthase α subunit contains the 3Fe-4S cluster of glutamate synthase, as shown by low-temperature EPR spectroscopy experiments. The glutamate synthase α subunit catalyzes the synthesis of glutamate from L- glutamine

and 2-oxoglutarate, provided that a reducing system (dithionite and methyl viologen) is present. The FMN moiety but not the 3Fe-4S cluster of the subunit appears to participate in this reaction. Furthermore, the isolated α subunit of glutamate synthase exhibits a glutaminase activity, which is absent in the glutamate synthase holoenzyme. These findings support a model for glutamate synthase according to which the enzymes prepared from various sources share a common glutamate synthase function (the α subunit of the bacterial enzyme, or its homologous polypeptide forming the ferredoxin-dependent plant enzyme) but differ for the chosen electron donor. The pyridine nucleotide-dependent forms of the enzyme have recruited a FAD-dependent oxidoreductase (the bacterial β subunit) to mediate electron transfer from the NAD(P)H substrate to the glutamate synthase polypeptide. However, it appears that the presence of the enzyme β subunit and/or of the additional iron-sulfur clusters (Centers II and III) of the bacterial glutamate synthase is required for communication between Center I (the 3Fe-4S center) and the FMN moiety within the α subunit, and for ensuring coupling of glutamine hydrolysis to the transfer of the released ammonia molecule to 2-oxoglutarate in the holoenzyme.

L6 ANSWER 95 OF 122 CAPLUS COPYRIGHT 2006 ACS on STN DUPLICATE 41

ACCESSION NUMBER: 1993:403879 CAPLUS

DOCUMENT NUMBER: 119:3879

TITLE: Substitution of Glu841 by lysine in the carbamate domain of carbamyl phosphate synthetase alters the catalytic properties of the glutaminase subunit

AUTHOR(S): Lusty, Carol J.; Liao, May

CORPORATE SOURCE: Dep. Mol. Genet., Public Health Res. Inst., New York, NY, 10016, USA

SOURCE: Biochemistry (1993), 32(5), 1278-84

CODEN: BICHAW; ISSN: 0006-2960

DOCUMENT TYPE: Journal

LANGUAGE: English

AB In previous studies a Glu841 \rightarrow Lys replacement in the carbamate phosphorylating domain located in the COOH half of the synthetase subunit of *Escherichia coli* carbamyl phosphate synthetase was shown to reduce overall synthesis of carbamyl phosphate by 4 orders of magnitude with either glutamine or NH_3 as nitrogen donor (Guillou, F.; et al., 1992). In the present study, the mutant enzyme has been further analyzed for its glutamine hydrolytic activity. The glutaminase activity of the mutant enzyme has the following properties. (1) In the absence of other substrates the turnover number is only marginally different from that of the wild-type complex. (2) The K_m for glutamine is 60 times higher than in wild-type complex and three times higher than in the separated glutaminase subunit. (3) In the present study wild-type carbamyl phosphate synthetase has been shown to catalyze glutamine hydrolysis by a mechanism involving an enzyme-bound acyl ester intermediate (γ -glutamyl thioester). This intermediate is formed and is hydrolyzed with rates consistent with overall glutamine hydrolysis. At physiol. concns. of glutamine (1.2 mM), the steady-state concentration of γ -glutamyl thioester is 0.3 mol/mol of wild-type enzyme. Under the same conditions, only 0.02 mol of thioester is measured in the mutant enzyme. Maximal accumulation of this covalent intermediate by the mutant enzyme required 10 times higher concns. of free glutamine. (4) The rate of reaction with 2-amino-4-oxo-5-chloropentanoate, a glutamine analog known to specifically alkylate an active site cysteine residue, is 2 orders of magnitude slower in the mutant. (5) Binding of both MgATP and bicarbonate to carbamylphosphate synthetase normally stimulates glutamine hydrolysis by 100-fold. This activation, presumed to be dependent on a carboxyphosphate-induced conformational change of the glutaminase active site, is not observed with the Lys841 enzyme. (6) Finally, the pH dependence of the glutaminase activity in the mutant complex is identical to that of the separated glutaminase subunit which exhibits fewer titratable groups than wild-type holoenzyme. Most of the properties listed above are also displayed by the

isolated glutaminase subunit. In addition to the previously reported effects on catalytic activity of the synthetase component, the Lys841 substitution therefore appears to uncouple functional interactions between the glutaminase and carbamate phosphorylation domains.

L6 ANSWER 102 OF 122 CAPLUS COPYRIGHT 2006 ACS on STN
ACCESSION NUMBER: 1993:142405 CAPLUS
DOCUMENT NUMBER: 118:142405
TITLE: Mechanistic studies of glutaminase activity
of a glutamine amidotransferase, carbamoyl
phosphate synthetase from Escherichia
coli
AUTHOR(S): Chang, Sun Hee Kim
CORPORATE SOURCE: Texas A and M Univ., College Station, TX, USA
SOURCE: (1991) 131 pp. Avail.: Univ. Microfilms Int., Order
No. DA9206471
From: Diss. Abstr. Int. B 1992, 52(12, Pt. 1), 6361-2
DOCUMENT TYPE: Dissertation
LANGUAGE: English
AB Unavailable

L6 ANSWER 112 OF 122 BIOTECHNO COPYRIGHT 2006 Elsevier Science B.V. on STN
DUPLICATE
ACCESSION NUMBER: 1985:15064463 BIOTECHNO
TITLE: The gene coding for carbamoyl-phosphate synthetase I
was formed by fusion of an ancestral glutaminase gene
and a synthetase gene
AUTHOR: Nyunoya H.; Broglie K.E.; Lusty C.J.
CORPORATE SOURCE: Molecular Genetics Laboratory, The Public Health
Research Institute of The City of New York, Inc., New
York, NY 10016, United States.
SOURCE: Proceedings of the National Academy of Sciences of the
United States of America, (1985), 82/8 (2244-2246)
CODEN: PNASA6
DOCUMENT TYPE: Journal; Article
COUNTRY: United States
LANGUAGE: English

AN 1985:15064463 BIOTECHNO
AB A near full-length cDNA copy of rat carbamoyl-phosphate
synthetase I (EC 6.3.4.16) mRNA has been cloned. The cDNA insert
in the recombinant plasmid pHN234 is 5.3 kilobases long. Analysis of the
sequence coding for carbamoyl-phosphate synthetase I indicates
that the gene has arisen from a fusion of two ancestral genes: one
homologous to Escherichia coli carA, coding for a
glutaminase subunit, and the second homologous to the carB gene
that codes for the synthetase subunit. A short amino acid
sequence previously proposed to be part of the active site involved in
glutamine amide nitrogen transfer in the E. coli and
yeast carbamoyl-phosphate synthetases (EC 6.3.5.5) is also
present in the rat enzyme. In the mammalian enzyme, however, the
glutaminase domain lacks a cysteine residue previously shown to
interact with glutamine. The cysteine is replaced by a serine
residue. This substitution could, in part, account for the inability of
mammalian carbamoyl-phosphate synthetase I to catalyze the
hydrolysis of glutamine to glutamic acid and ammonia.

L6 ANSWER 122 OF 122 CAPLUS COPYRIGHT 2006 ACS on STN
ACCESSION NUMBER: 1967:18107 CAPLUS
DOCUMENT NUMBER: 66:18107
TITLE: Bacterial production of glutamic acid in stored
comminuted beef
AUTHOR(S): Gardner, G. A.; Stewart, David John
CORPORATE SOURCE: Queen's Univ., Belfast, Ire.
SOURCE: Journal of Applied Bacteriology (1966), 29(2), 365-74
CODEN: JABAA4; ISSN: 0021-8847

DOCUMENT TYPE: Journal
LANGUAGE: English

AB The production of free glutamic acid from the deamidation of glutamine in stored meat was due to bacterial activity and not to glutaminase in the meat. Pseudomonas-Achromobacter species predominated after 41 hrs. at 15°. The glutaminase of an isolated pseudomonad was optically active at 36° and pH 5, and was constitutive.

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 USPATFULL, ESBIOBASE, BIOTECHNO, LIFESCI' ENTERED AT 16:33:51 ON 28 AUG
 2006

L2 1535 SEA GLUTAMINAS? AND GLUTAMINE? AND (CORYNEFOR? OR GLUTAMICUM?
 OR COLI? OR BACTER? OR BREVIBACT?)
 L3 1174 SEA L2 AND (METHOD? OR PRODUCT? OR SYNTH?)
 L4 342 SEA GLUTAMINAS?(S) (GLUTAMINE?) (S) (CORYNEFOR? OR GLUTAMICUM? OR
 COLI? OR BACTER? OR BREVIBACT?)
 L5 211 SEA L4(S) (METHOD? OR SYNTH? OR PRODUCT?)
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 FILE LAST UPDATED: 26 Aug 2006 (20060826/UP). FILE COVERS 1950 TO DATE.

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